

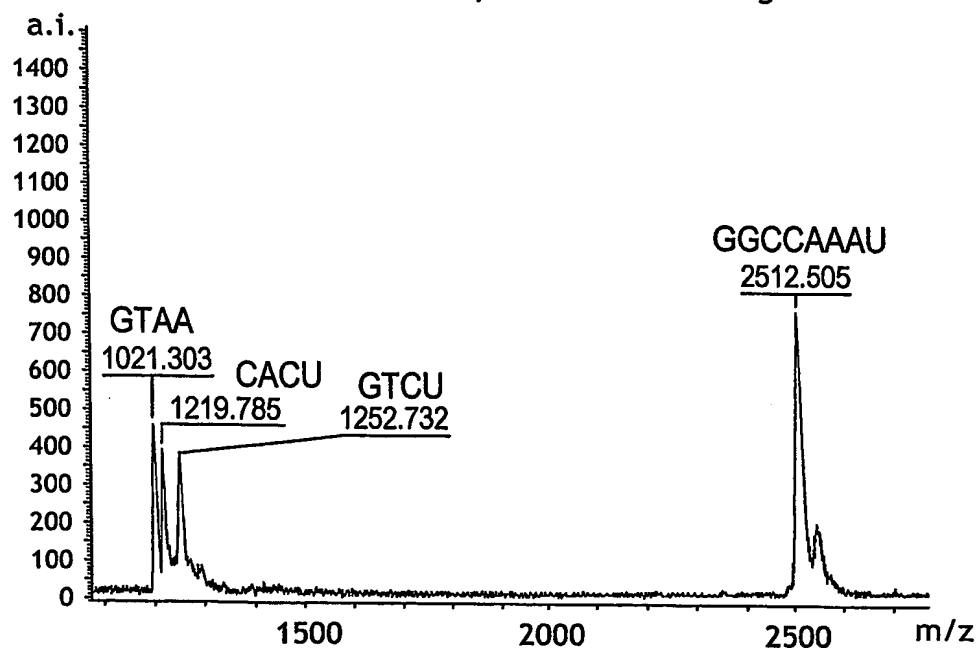
1 / 8

FIG.1

5' -GTCU|CTU|CACU|U|GGCCAAAU|GTAA

Fragment	Expected Mass	Found Mass
GTCUCTUCACUUGGCCAAAU GTAA	7312,9	
CTU	918,6	
GTAA	1197,9	1201,3
CACU	1216,8	1219,8
GTCU	1247,8	1252,7
GGCCAAAU	2501,7	2512,5

Clivage par NH₄OH de GTCUCTUCACUUGGCCAAAU GTAA
Matrice HPA mode positif sur anchortaget400



2 / 8

FIG.2

5' -GTC|TC|TTCAC|TTGGC|C|AAATGTAA

Fragment	Expected Mass	Found Mass
GTCTCTTCACTTGGCCAAATGTAA	7398,9	
AC	637,4	
TTC	932,6	928,5
GTC	957,6	953,6
TTGGC	1591,0	1588,2
AAATGTA	2128,5	2127
AAATGTAA	2441,8	2440,5

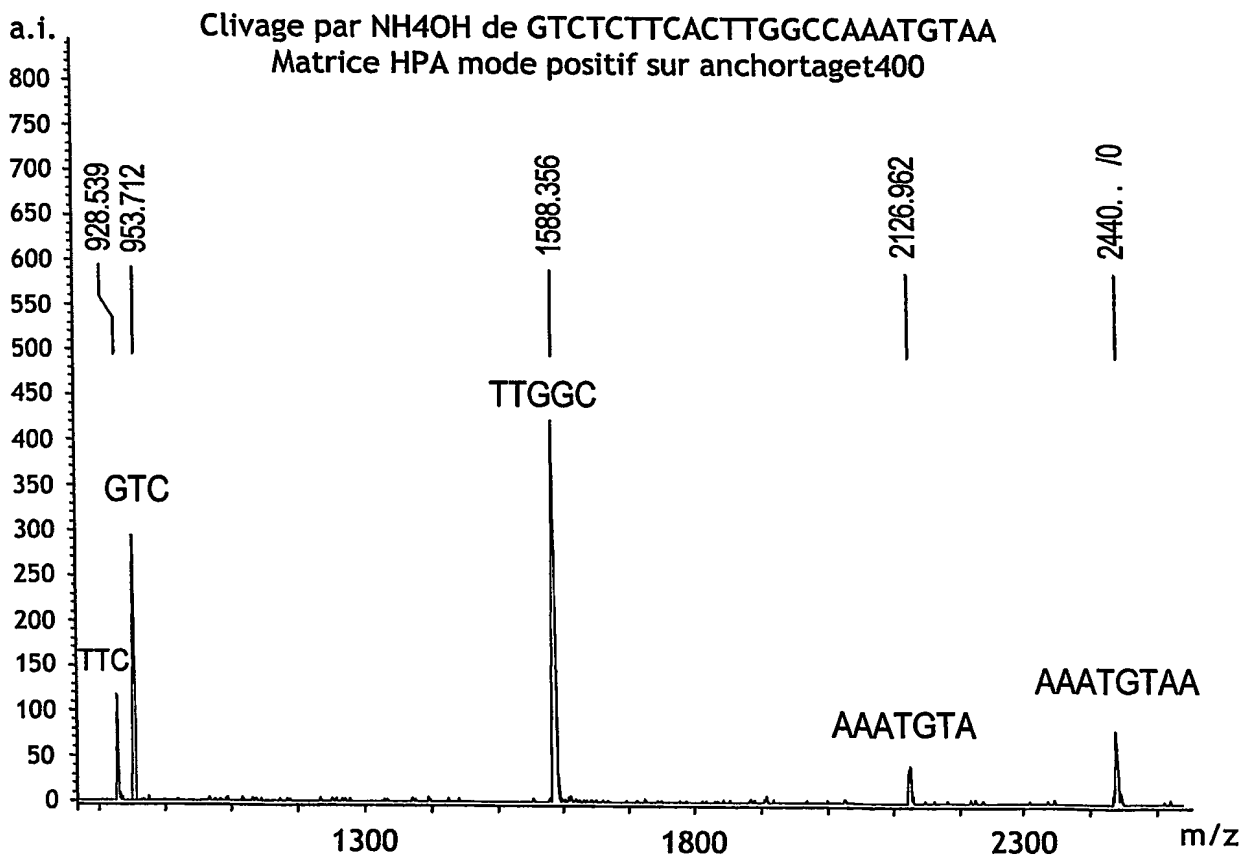


Figure 3

Fragments at 100% Exchange dNTP for NTP

5'-GTCTCTT CACTT GGCCAAATGTAA
5'-GTCCTTCAC TTGGCCAAATGTAA
5'-GTCTGTTCA CTGGCCA AATGTAA
5'-GTCCTTCAC TTG G CAAATGTAA

Additional Fragments at 70% Exchange dNTP for NTP

5'-GTCTCTTCACTTGGCCAAATGTAA (25%)
5'-GTCTCTTCACTTGGCCAAATGTAA (25%)

5'-GTC/TCTTC/ACTTGGCC/AAATGTAA (25%)
5'-GTCTCTTCAC/TTGGCC/AAATGTAA (25%)

5'-GTCTCTTCACTTGGCCAAATGTAA (25%)
5'-GTCTCTTCACTTGGCCAAATGTAA (25%)

5'-GTTCTCTTCAGTTTCGCCAAATGTA (25%)
5'-GTTCTCTTCAGTTTCGCCAAATGTA (25%)

Figure 4

basic sequence:

5'-TTCACCTGGCCAAATGT^{RNA}AAGNGAAGAACAGAGTC-3'

complementary template sequences:

3'-AAGTGAACCGGTTTACATTCCTTCTTGTCTCAG-5' G template

3'-AAGTGAACCGGTTTACATTCGCTTCTTGTCTCAG-5' C template

3'-AAGTGAACCGGTTTACATTCTCTTCTTGTCTCAG-5' A template

primer sequence:

5'-TTCACCTGGCCAAATGT^{RNA}AAG-3'

sequences after extension and before cleavage:

5'-TTCACCTGGCCAAATGT^{RNA}AAGG^{RNA}G^{RNA}AAG^{RNA}AACAG^{RNA}AG^{RNA}TC-3' N = G

5'-TTCACCTGGCCAAATGT^{RNA}AAGCG^{RNA}AAG^{RNA}AACAG^{RNA}AG^{RNA}TC-3' N = C

5'-TTCACCTGGCCAAATGT^{RNA}AAGAG^{RNA}AAG^{RNA}AACAG^{RNA}AG^{RNA}TC-3' N = A

Fragments after cleavage:

AAGG^{RNA} G^{RNA} AAG^{RNA} AACAG^{RNA} AG^{RNA} TC Fragments for N = G

AAGCG^{RNA} AAG^{RNA} AACAG^{RNA} AG^{RNA} TC Fragments for N = C

AAGAG^{RNA} AAG^{RNA} AACAG^{RNA} AG^{RNA} TC Fragments for N = A

Signals that differ:

AAGG^{RNA} for N = G = 1319 Da

AAGCG^{RNA} for N = C = 1633 Da

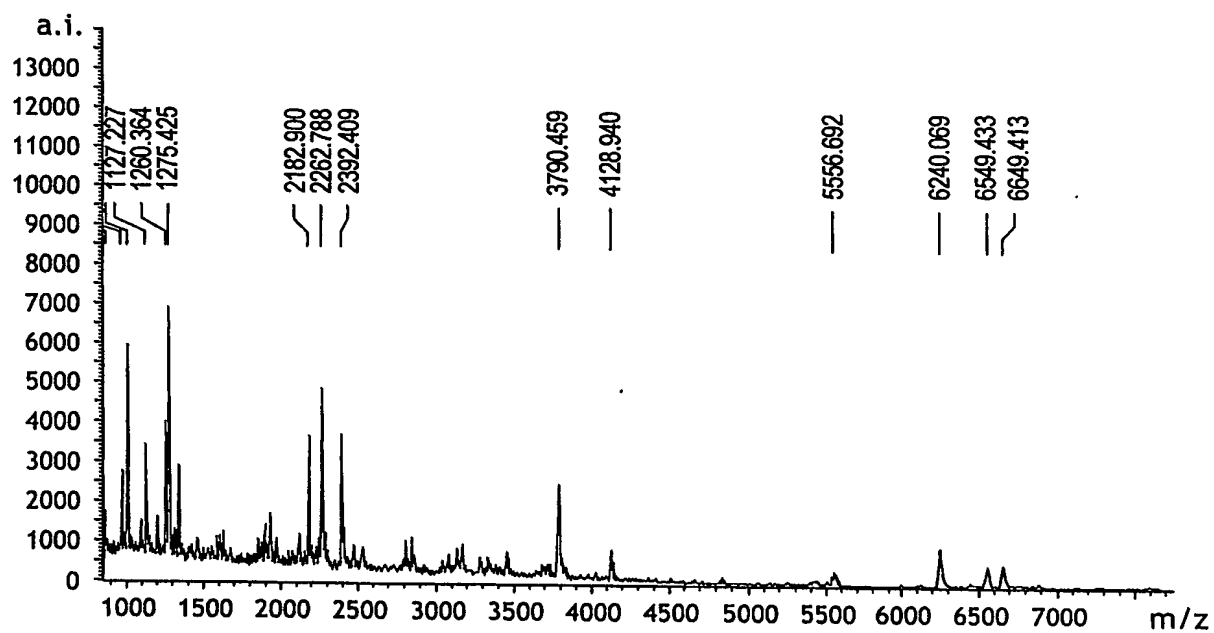
AAGAG^{RNA} for N = A = 1609 Da

Signals that are the same:

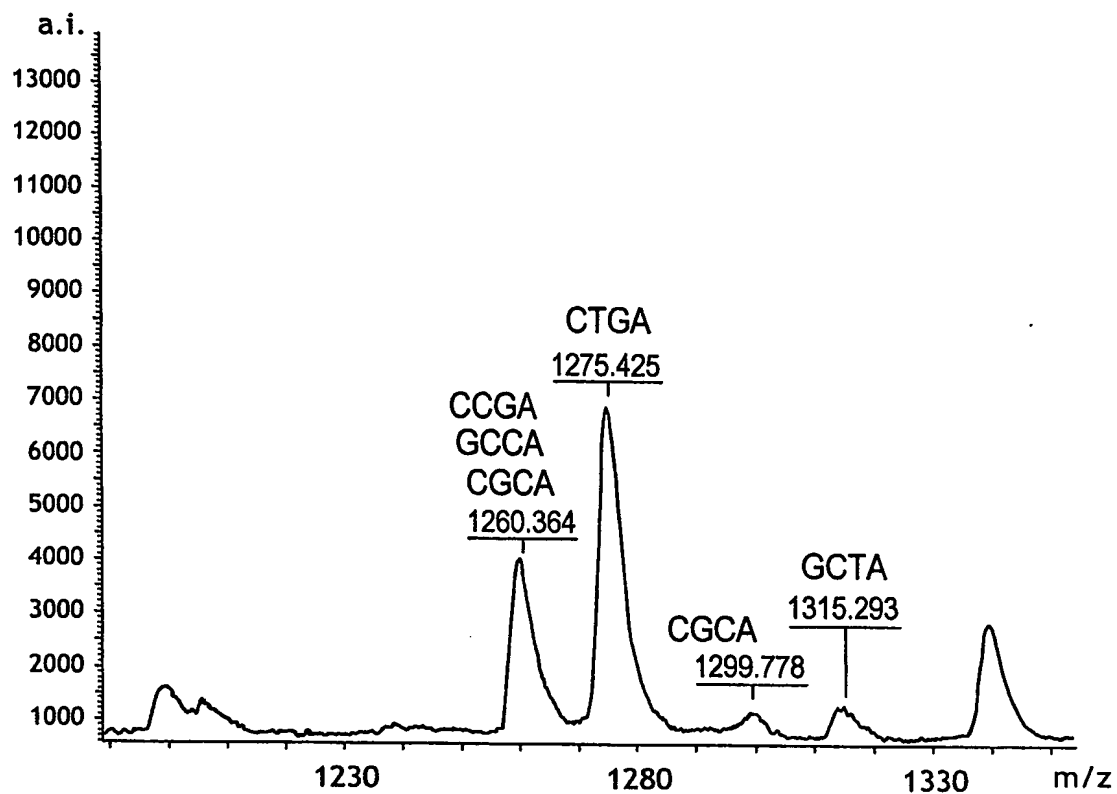
AACAG^{RNA} = 1593 Da

AAG^{RNA} = 990 Da

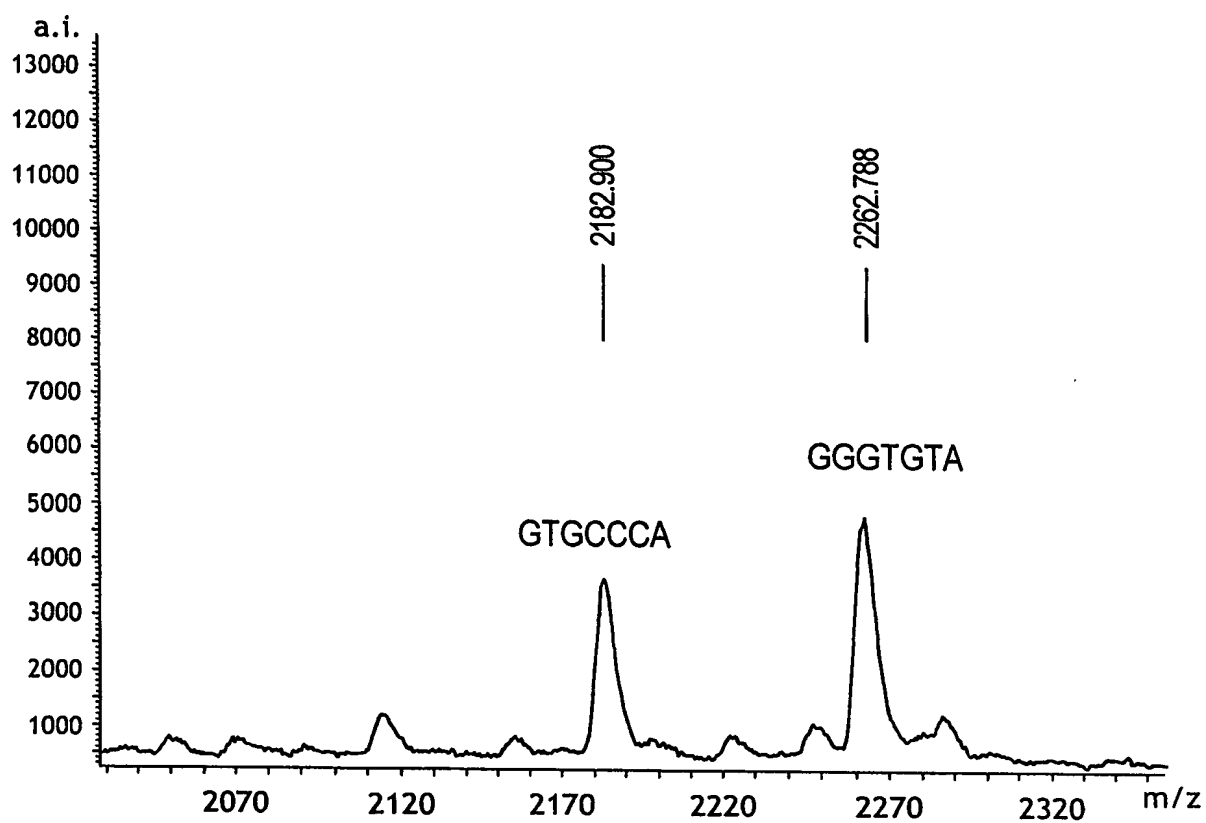
5 / 8

FIG.5

6 / 8

FIG.6

7/8

FIG.7

8 / 8

FIG.8

PCR 216 bp Fragment Forward Atp/dAtp

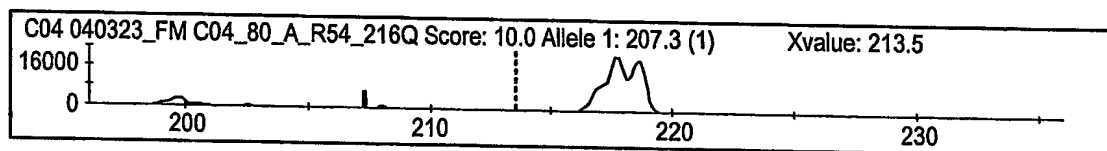
67% Atp/dAtp



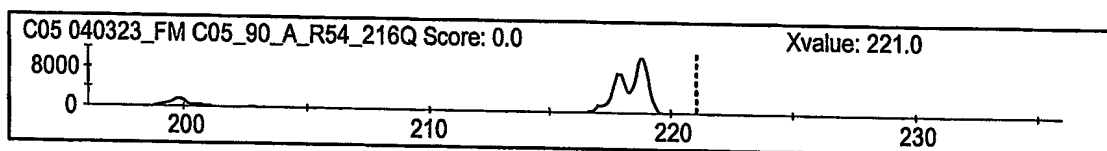
75% Atp/dAtp



80% Atp/dAtp



90% Atp/dAtp



100% Atp

